atg Met 1

ctg Leu

ttc Phe

cgt Arg

SEQUENCE LISTING

<110>	Deisher, Theresa A. Conklin, Darrell C. Raymond, Fenella Bukowski, Thomas R. Holderman, Susan D. Hansen, Birgit Sheppard, Paul O.	
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	ctg cgg ctg tac cag ctc tac agc cgg acc agt ggg a Leu Arg Leu Tyr Gln Leu Tyr Ser Arg Thr Ser Gly L 55 60	

,,

		_	-	_		-			Ser	-	-			gat Asp		240
-	_		_	-			-			-				agt Ser 95		288
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		_				_		_			_			tgt Cys		384
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	gtg Val												Pro	gcc Ala		621
cate tgae aace	caga caaaa agaca	gga a aga d aac 1	atat ctca tcta	tttt cgca aact	ac a aa g cg t	tgaa ggac cccc	aaata tgtag agagg	a agg g tca g agg	gatt aacc gact	ttat caca tgaa	tgt ggt tga	tgac gctt ggaa	ttg gtc acc	aaac tctc	aaactg ccccga tctagg ctttga ag	68: 74: 80: 86: 91:

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Phe Arg Ile His Val Glu Asn Gln Thr Arg Ala Arg Asp Asp Val Ser
Arg Lys Gln Leu Arg Leu Tyr Gln Leu Tyr Ser Arg Thr Ser Gly Lys
His Ile Gln Val Leu Gly Arg Arg Ile Ser Ala Arg Gly Glu Asp Gly
Asp Lys Tyr Ala Gln Leu Leu Val Glu Thr Asp Thr Phe Gly Ser Gln
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Val Arg Ile Lys Gly Lys Glu Thr Glu Phe Tyr Leu Cys Met Asn Arg
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Lys Gly Lys Leu Val Gly Lys Pro Asp Gly Thr Ser Lys Glu Cys Val
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Phe Ile Glu Lys Val Leu Glu Asn Asn Tyr Thr Ala Leu Met Ser Ala
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Lys Tyr Ser Gly Trp Tyr Val Gly Phe Thr Lys Lys Gly Arg Pro Arg
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Lys Gly Pro Lys Thr Arg Glu Asn Gln Gln Asp Val His Phe Met Lys
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Arg Tyr Pro Lys Gly Gln Pro Glu Leu Gln Lys Pro Phe Lys Tyr Thr
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		220> 223>	5' 1	linke	er se	equer	nce								-	•
cgca	nttgo ntcca	acg t	ctaaa	gaaco	a ga		-	-	_	_					jacttc tgcgg	60 120 141
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Pro	Ser		20 Asp	Gly	Arg	Ser		25 Cys	Glu	Arg	His	Va 1 45	30 Leu	Gly	Va1	
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Arg 65		Glu	Pro	Gln	Leu 70	55 Lys	Gly	Ile	Val	Thr 75	60 Arg	Leu	Phe	Ser	Gln 80	
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Leu Arg Val Val Ala Ile Gln Gly Val Lys Ala Ser Leu Tyr Val Ala
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                            120
                                                 125
Met Asn Gly Glu Gly Tyr Leu Tyr Ser Ser Asp Val Phe Thr Pro Glu
                        135
Cys Lys Phe Lys Glu Ser Val Phe Glu Asn Tyr Tyr Val Ile Tyr Ser
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Ser Thr Leu Tyr Arg Gln Gln Glu Ser Gly Arg Ala Trp Phe Leu Gly
                                    170
                165
Leu Asn Lys Glu Gly Gln Ile Met Lys Gly Asn Arg Val Lys Lys Thr
Lys Pro Ser Ser His Phe Val Pro Lys Pro Ile Glu Val Cys Met Tyr
                            200
Arg Glu Pro Ser Leu His Glu Ile Gly Glu Lys Gln Gly Arg Ser Arg
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Lys Ser Ser Gly Thr Pro Thr Met Asn Gly Gly Lys Val Val Asn Gln
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Asp Ser Thr
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Met Tyr Arg Glu Pro Ser Leu His Glu Ile Gly Glu Asn Lys Gly Val 145 150 155 160 Gln Gly Lys Phe Trp Thr Pro Pro 165

<210> 23

<211> 247

<212> PRT

<213> Homo sapiens

<400> 23

Met Ala Ala Ala Ile Ala Ser Gly Leu Ile Arg Gln Lys Arg Gln Ala Arg Glu Gln His Trp Asp Arg Pro Ser Ala Ser Arg Arg Arg Ser Ser Pro Ser Lys Asn Arg Gly Leu Cys Asn Gly Asn Leu Val Asp Ile Phe Ser Lys Val Arg Ile Phe Gly Leu Lys Lys Arg Arg Leu Arg Arg Gln Asp Pro Gln Leu Lys Gly Ile Val Thr Arg Leu Tyr Cys Arg Gln Gly 70 75 Tyr Tyr Leu Gln Met His Pro Asp Gly Ala Leu Asp Gly Thr Lys Asp Asp Ser Thr Asn Ser Thr Leu Phe Asn Leu Ile Pro Val Gly Leu Arg 105 Val Val Ala Ile Gln Gly Val Lys Thr Gly Leu Tyr Ile Ala Met Asn 120 Gly Glu Gly Tyr Leu Tyr Pro Ser Glu Leu Phe Thr Pro Glu Cys Lys -...135 140 Phe Lys Glu Ser Val Phe Glu Asn Tyr Tyr Val Ile Tyr Ser Ser Met Leu Tyr Arg Gln Gln Glu Ser Gly Arg Ala Trp Phe Leu Gly Leu Asn 165 170 Lys Glu Gly Gln Ala Met Lys Gly Asn Arg Val Lys Lys Thr Lys Pro 180 185 Ala Ala His Phe Leu Pro Lys Pro Leu Glu Val Ala Met Tyr Arg Glu 195 200 205 Pro Ser Leu His Asp Val Gly Glu Thr Val Pro Lys Pro Gly Val Thr 215 220 Pro Ser Lys Ser Thr Ser Ala Ser Ala Ile Met Asn Gly Gly Lys Pro 240 230 235 Val Asn Lys Ser Lys Thr Thr

<211> 245 <212> PRT <213> Homo sapiens

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Met Ala Ala Ala Ile Ala Ser Ser Leu Ile Arg Gln Lys Arg Gln Ala Arg Glu Arg Glu Lys Ser Asn Ala Cys Lys Cys Val Ser Ser Pro Ser Lys Gly Lys Thr Ser Cys Asp Lys Asn Lys Leu Asn Val Phe Ser Arg 40 Val Lys Leu Phe Gly Ser Lys Lys Arg Arg Arg Arg Pro Glu Pro Gln Leu Lys Gly Ile Val Thr Lys Leu Tyr Ser Arg Gln Gly Tyr His 70 75 Leu Gln Leu Gln Ala Asp Gly Thr Ile Asp Gly Thr Lys Asp Glu Asp Ser Thr Tyr Thr Leu Phe Asn Leu Ile Pro Val Gly Leu Arg Val Val Ala Ile Gln Gly Val Gln Thr Lys Leu Tyr Leu Ala Met Asn Ser Glu 120 Gly Tyr Leu Tyr Thr Ser Glu Leu Phe Thr Pro Glu Cys Lys Phe Lys 135 Glu Ser Val Phe Glu Asn Tyr Tyr Val Thr Tyr Ser Ser Met Ile Tyr 150 155 Arg Gln Gln Ser Gly Arg Gly Trp Tyr Leu Gly Leu Asn Lys Glu 170 165 Gly Glu Ile Met Lys Gly Asn His Val Lys Lys Asn Lys Pro Ala Ala 185 His Phe Leu Pro Lys Pro Leu Lys Val Ala Met Tyr Lys Glu Pro Ser Leu His Asp Leu Thr Glu Phe Ser Arg Ser Gly Ser Gly Thr Pro Thr 215 Lys Ser Arg Ser Val Ser Gly Val Leu Asn Gly Gly Lys Ser Met Ser 230 240 235 His Asn Glu Ser Thr 245

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<213> Homo sapiens

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Glu Pro Gly Gly Ser Arg Pro Val Ser Ala Gln Arg Arg Val Cys Pro
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Arg Gly Thr Lys Ser Leu Cys Gln Lys Gln Leu Leu Ile Leu Leu Ser
Lys Val Arg Leu Cys Gly Gly Arg Pro Ala Arg Pro Asp Arg Gly Pro
Glu Pro Gln Leu Lys Gly Ile Val Thr Lys Leu Phe Cys Arg Gln Gly
                                         75
Phe Tyr Leu Gln Ala Asn Pro Asp Gly Ser Ile Gln Gly Thr Pro Glu
Asp Thr Ser Ser Phe Thr His Phe Asn Leu Ile Pro Val Gly Leu Arg
                                105
Val Val Thr Ile Gln Ser Ala Lys Leu Gly His Tyr Met Ala Met Asn
Ala Glu Gly Leu Leu Tyr Ser Ser Pro His Phe Thr Ala Glu Cys Arg
                        135
                                             140
Phe Lys Glu Cys Val Phe Glu Asn Tyr Tyr Val Leu Tyr Ala Ser Ala
                                         155
                    150
Leu Tyr Arg Gln Arg Arg Ser Gly Arg Ala Trp Tyr Leu Gly Leu Asp
                165
                                    170
Lys Glu Gly Gln Val Met Lys Gly Asn Arg Val Lys Lys Thr Lys Ala
                                185
Ala Ala His Phe Leu Pro Lys Leu Leu Glu Val Ala Met Tyr Gln Glu
                            200
                                                 205
Pro Ser Leu His Ser Val Pro Glu Ala Ser Pro Ser Ser Pro Pro Ala
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                        215
                                             220
Pro
225
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Met Ser Gly Pro Gly Thr Ala Ala Val Ala Leu Leu Pro Ala Val Leu

1 5 10 15

Leu Ala Leu Leu Ala Pro Trp Ala Gly Arg Gly Gly Ala Ala Ala Pro
20 25 30

Thr Ala Pro Asn Gly Thr Leu Glu Ala Glu Leu Glu Arg Arg Trp Glu
35 40 45

Ser Leu Val Ala Leu Ser Leu Ala Arg Leu Pro Val Ala Ala Gln Pro
50 55 60

**

ri nrr

Lys Glu Ala Ala Val Gln Ser Gly Ala Gly Asp Tyr Leu Leu Gly Ile Lys Arg Leu Arg Arg Leu Tyr Cys Asn Val Gly Ile Gly Phe His Leu 90 Gln Ala Leu Pro Asp Gly Arg Ile Gly Gly Ala His Ala Asp Thr Arg 105 Asp Ser Leu Leu Glu Leu Ser Pro Val Glu Arg Gly Val Val Ser Ile 120 Phe Gly Val Ala Ser Arg Phe Phe Val Ala Met Ser Ser Lys Gly Lys 135 Leu Tyr Gly Ser Pro Phe Phe Thr Asp Glu Cys Thr Phe Lys Glu Ile 155 Leu Leu Pro Asn Asn Tyr Asn Ala Tyr Glu Ser Tyr Lys Tyr Pro Gly 170 165 Met Phe Ile Ala Leu Ser Lys Asn Gly Lys Thr Lys Lys Gly Asn Arg 185 Val Ser Pro Thr Met Lys Val Thr His Phe Leu Pro Arg Leu 195 200

<210> 27

<211> 208

<212> PRT

<213> Homo sapiens

<400> 27

Met Ala Leu Gly Gln Lys Leu Phe Ile Thr Met Ser Arg Gly Ala Gly Arg Leu Gln Gly Thr Leu Trp Ala Leu Val Phe Leu Gly Ile Leu Val Gly Met Val Val Pro Ser Pro Ala Gly Thr Arg Ala Asn Asn Thr Leu Leu Asp Ser Arg Gly Trp Gly Thr Leu Leu Ser Arg Ser Arg Ala Gly 55 Leu Ala Gly Glu Ile Ala Gly Val Asn Trp Glu Ser Gly Tyr Leu Val Gly Ile Lys Arg Gln Arg Arg Leu Tyr Cys Asn Val Gly Ile Gly Phe 85 90 His Leu Gln Val Leu Pro Asp Gly Arg Ile Ser Gly Thr His Glu Glu Asn Pro Tyr Ser Leu Leu Glu Ile Ser Thr Val Glu Arg Gly Val Val 120 Ser Leu Phe Gly Val Arg Ser Ala Leu Phe Val Ala Met Asn Ser Lys 135 140 Gly Arg Leu Tyr Ala Thr Pro Ser Phe Gln Glu Glu Cys Lys Phe Arg **~** 150 155

101 2 At 0.0

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Glu Thr Leu Leu Pro Asn Asn Tyr Asn Ala Tyr Glu Ser Asp Leu Tyr
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                                    170
Gln Gly Thr Tyr Ile Ala Leu Ser Lys Tyr Gly Arg Val Lys Arg Gly
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Ser Lys Val Ser Pro Ile Met Thr Val Thr His Phe Leu Pro Arg Ile
                            200
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Gly Ser Gly Ala Phe Pro Pro Gly His Phe Lys Asp Pro Lys Arg Leu
Tyr Cys Lys Asn Gly Gly Phe Phe Leu Arg Ile His Pro Asp Gly Arg
Val Asp Gly Val Arg Glu Lys Ser Asp Pro His Ile Lys Leu Gln Leu
Gln Ala Glu Glu Arg Gly Val Val Ser Ile Lys Gly Val Cys Ala Asn
Arg Tyr Leu Ala Met Lys Glu Asp Gly Arg Leu Leu Ala Ser Lys Cys
                                    90
Val Thr Asp Glu Cys Phe Phe Phe Glu Arg Leu Glu Ser Asn Asn Tyr
Asn Thr Tyr Arg Ser Arg Lys Tyr Thr Ser Trp Tyr Val Ala Leu Lys
                            120
Arg Thr Gly Gln Tyr Lys Leu Gly Ser Lys Thr Gly Pro Gly Gln Lys
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Ala Ile Leu Phe Leu Pro Met Ser Ala Lys Ser
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Thr Arg Asp Arg Ser Asp Gln His Ile Gln Leu Gln Leu Ser Ala Glu
                       55
Ser Val Gly Glu Val Tyr Ile Lys Ser Thr Glu Thr Gly Gln Tyr Leu
Ala Met Asp Thr Asp Gly Leu Leu Tyr Gly Ser Gln Thr Pro Asn Glu
                                    90
Glu Cys Leu Phe Leu Glu Arg Leu Glu Glu Asn His Tyr Asn Thr Tyr
                                105
Ile Ser Lys Lys His Ala Glu Lys Asn Trp Phe Val Gly Leu Lys Lys
       115
                            120
Asn Gly Ser Cys Lys Arg Gly Pro Arg Thr His Tyr Gly Gln Lys Ala
                       135
Ile Leu Phe Leu Pro Leu Pro Val Ser Ser Asp
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                    150
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Pro Gly Cys Cys Cys Cys Phe Leu Leu Phe Leu Val Ser Ser
Val Pro Val Thr Cys Gln Ala Leu Gly Gln Asp Met Val Ser Pro Glu
Ala Thr Asn Ser Ser Ser Ser Phe Ser Ser Pro Ser Ser Ala Gly
Arg His Val Arg Ser Tyr Asn His Leu Gln Gly Asp Val Arg Trp Arg
Lys Leu Phe Ser Phe Thr Lys Tyr Phe Leu Lys Ile Glu Lys Asn Gly
Lys Val Ser Gly Thr Lys Lys Glu Asn Cys Pro Tyr Ser Ile Leu Glu
                                105
Ile Thr Ser Val Glu Ile Gly Val Val Ala Val Lys Ala Ile Asn Ser
                            120
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Tyr Asn Thr Tyr Ala Ser Phe Asn Trp Gln His Asn Gly Arg Gln Met 170 175

Asn Tyr Tyr Leu Ala Met Asn Lys Lys Gly Lys Leu Tyr Gly Ser Lys

Glu Phe Asn Asn Asp Cys Lys Leu Lys Glu Arg Ile Glu Glu Asn Gly

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150

165

155

Tyr Val Ala Leu Asn Gly Lys Gly Ala Pro Arg Arg Gly Gln Lys Thr Arg Arg Lys Asn Thr Ser Ala His Phe Leu Pro Met Val Val His Ser 205 195 200

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<211> 194

<212> PRT

<213> Homo sapiens

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Lys Lys Thr Lys Lys Glu Gln Lys Thr Ala His Phe Leu Pro Met Ala

185

170

190

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Ile Thr

<211> 233

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<213> Homo sapiens

180

165

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Val Ser Gln Gln His Val Arg Glu Gln Ser Leu Val Thr Asp Gln Leu
Ser Arg Arg Leu Ile Arg Thr Tyr Gln Leu Tyr Ser Arg Thr Ser Gly
Lys His Val Gln Val Leu Ala Asn Lys Arg Ile Asn Ala Met Ala Glu
Asp Gly Asp Pro Phe Ala Lys Leu Ile Val Glu Thr Asp Thr Phe Gly
                                 105
Ser Arg Val Arg Val Arg Gly Ala Glu Thr Gly Leu Tyr Ile Cys Met
Asn Lys Lys Gly Lys Leu Ile Ala Lys Ser Asn Gly Lys Gly Lys Asp
                         135
                                             140
Cys Val Phe Thr Glu Ile Val Leu Glu Asn Asn Tyr Thr Ala Leu Gln
Asn Ala Lys Tyr Glu Gly Trp Tyr Met Ala Phe Thr Arg Lys Gly Arg
                 165
                                     170
Pro Arg Lys Gly Ser Lys Thr Arg Gln His Gln Arg Glu Val His Phe
                                 185
Met Lys Arg Leu Pro Arg Gly His His Thr Thr Glu Gln Ser Leu Arg
                             200
Phe Glut Phe Leu Asn Tyr Pro Pro Phe Thr Arg Ser Leu Arg Gly Ser
                         215
                                             220
Gln Arg Thr Trp Ala Pro Glu Pro Arg
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                     230
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<210> 33

<211> 268

<212> PRT

<213> Homo sapiens

<400> 33

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Ser Ala Trp Ala His Gly Glu Lys Arg Leu Ala Pro Lys Gly Gln Pro 20 25 30

Gly Pro Ala Ala Thr Asp Arg Asn Pro Ile Gly Ser Ser Ser Arg Gln 35 40 45

Ser Ser Ser Ser Ser Ala Met Ser Ser Ser Ser Ala Ser Ser Ser Pro Ala 50 60

an arm of

Ala Ser Leu Gly Ser Gln Gly Ser Gly Leu Glu Gln Ser Ser Phe Gln Trp Ser Pro Ser Gly Arg Arg Thr Gly Ser Leu Tyr Cys Arg Val Gly 90 Ile Gly Phe His Leu Gln Ile Tyr Pro Asp Gly Lys Val Asn Gly Ser His Glu Ala Asn Met Leu Ser Val Leu Glu Ile Phe Ala Val Ser Gln 120 Gly Ile Val Gly Ile Arg Gly Val Phe Ser Asn Lys Phe Leu Ala Met 135 140 Ser Lys Lys Gly Lys Leu His Ala Ser Ala Lys Phe Thr Asp Asp Cys 150 Lys Phe Arg Glu Arg Phe Gln Glu Asn Ser Tyr Asn Thr Tyr Ala Ser 170 Ala Ile His Arg Thr Glu Lys Thr Gly Arg Glu Trp Tyr Val Ala Leu Asn Lys Arg Gly Lys Ala Lys Arg Gly Cys Ser Pro Arg Val Lys Pro 200 Gln His Ile Ser Thr His Phe Leu Pro Arg Phe Lys Gln Ser Glu Gln 215 Pro Glu Leu Ser Phe Thr Val Thr Val Pro Glu Lys Lys Asn Pro Pro 230 Ser Pro Ile Lys Ser Lys Ile Pro Leu Ser Ala Pro Arg Lys Asn Thr Asn Ser Val Lys Tyr Arg Leu Lys Phe Arg Phe Gly 260

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<213> Homo sapiens

<400> 34

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Val Pro Phe Gly Asn Val Pro Val Leu Pro Val Asp Ser Pro Val Leu 20 25 30
Leu Ser Asp His Leu Gly Gln Ser Glu Ala Gly Gly Leu Pro Arg Gly 35 40 45
Pro Ala Val Thr Asp Leu Asp His Leu Lys Gly Ile Leu Arg Arg 50 55 60
Gln Leu Tyr Cys Arg Thr Gly Phe His Leu Glu Ile Phe Pro Asn Gly 65 70 75 80
Thr Ile Gln Gly Thr Arg Lys Asp His Ser Arg Phe Gly Ile Leu Glu 85 90 95

Him district to the second

Phe Ile Ser Ile Ala Val Gly Leu Val Ser Ile Arg Gly Val Asp Ser Gly Leu Tyr Leu Gly Met Asn Glu Lys Gly Glu Leu Tyr Gly Ser Glu 120 Lys Leu Thr Gln Glu Cys Val Phe Arg Glu Gln Phe Glu Glu Asn Trp 135 Tyr Asn Thr Tyr Ser Ser Asn Leu Tyr Lys His Val Asp Thr Gly Arg 150 155 Arg Tyr Tyr Val Ala Leu Asn Lys Asp Gly Thr Pro Arg Glu Gly Thr 165 170 Arg Thr Lys Arg His Gln Lys Phe Thr His Phe Leu Pro Arg Pro Val 185 Asp Pro Asp Lys Val Pro Glu Leu Tyr Lys Asp Ile Leu Ser Gln Ser 200 <210> 35 <211> 239 <212> PRT

<213> Homo sapiens

<400> 35

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Arg Arg Arg Gln Lys Gln Ser Pro Asp Asn Leu Glu Pro Ser His
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                                            220
Val Gln Ala Ser Arg Leu Gly Ser Gln Leu Glu Ala Ser Ala His
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		_				tac Tyr 55							_		192
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aag (Lys (ggt o Gly F	ccc a Pro l	_ys ¯	acc d Thr # 165	gc g Arg G	jag a Nu <i>P</i>	iac (Isn (Gln (caa g G1n A 170	yat g Asp V	yta c /al H	ac t lis P	he M	tg a let L 75	ag ys	528
cgt 1 Arg	tac (Tyr I	Pro l	aag (_ys (180	gga (Gly (cag g Gln A	gcc g Na (alu I	ctg (Leu (185	cag a Gln l	aag (_ys f	ccc t Pro F	he L	iaa t .ys T .90	ac a yr T	icc Thr	576
aca (Val i	acc a Thr 1 195	aag Lys	cga t Arg :	tcc (Ser /	Arg /	egg Arg 200	atc Ile	cgc (Arg l	ccc a	ihr b	cac d dis F 205	cc (Pro (ggc t Ny	ag *	624
aagg tctt caca aact ccaa	aaga ggga aggg catc aggt	at c gg a ga c cc c tc t	tcta gggg cgct agag gaaa	tttt cgat gtca gagg gcaa	t gta a gga a cca a cta a ca	acat atto caca tgaa aaaa	tgtg cact ggtg cgag aaaa	ttt gtt ctt gaa aaa	aaaa gacc gcct actg	gaa tga ctc cga aaa	gacaa accca tcta gaaa	aaaa catg ggag ccaa	et ga ac aa gt ga ag ta	aacct aagga acaat cctt	aaaat taaag actca ttcaa tcccc aaaaa	684 744 804 864 924 984 1023
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